

Pixel Barrel Module Alignment Using Overlaps

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Introduction

- Develop alternative strategy that provides cross check and ultimately improve the pixel alignment.
- Constraining the neighboring modules together using overlaps.
- Overlap is defined for a track passing through two neighboring modules in the same layer and in the same eta ring.
- Advantage of this reduces a large number of modules into a small number of regions that requires fewer degrees of freedom to solve in the global χ^2 .

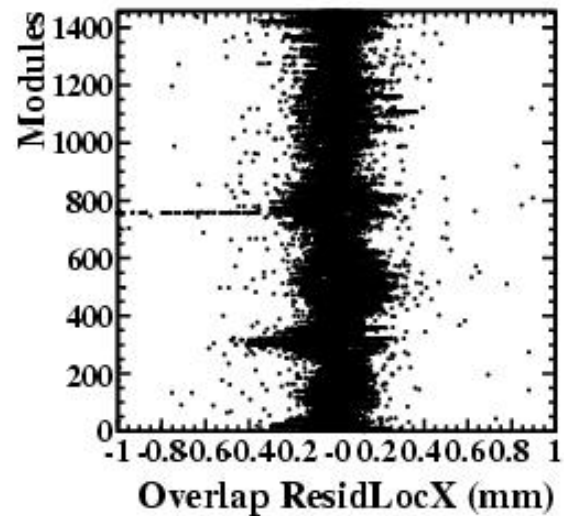
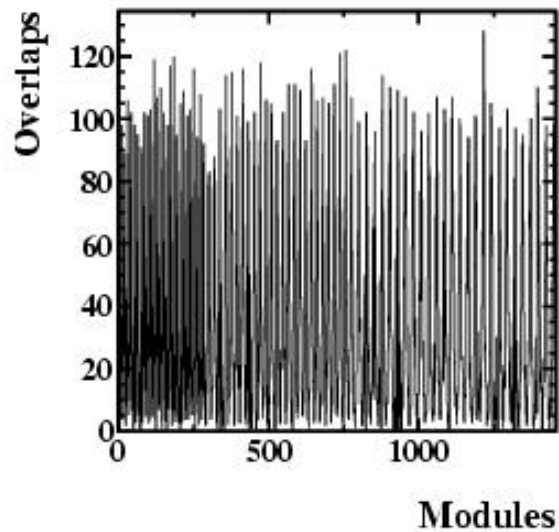
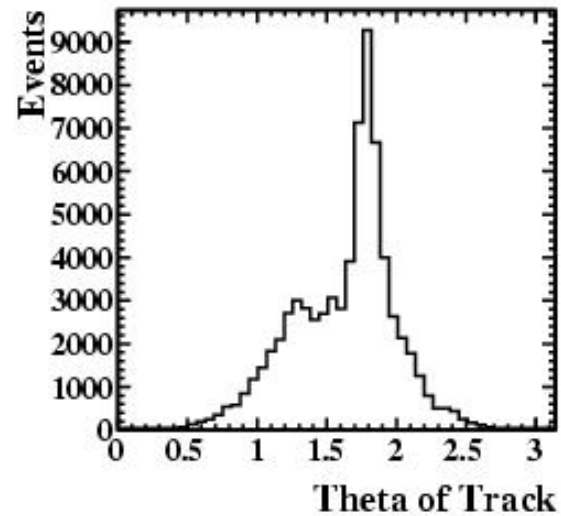
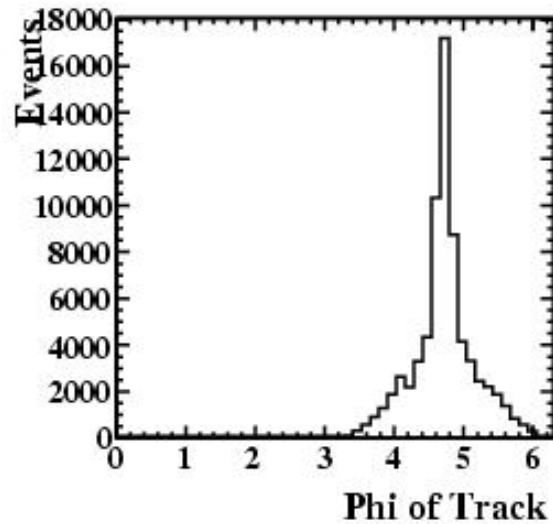


Datasets and Event Selections

- Using most of cosmic bfield off data from Max, about 180K.
- Reprocessed with default cosmic tracking with latest cosmic 03 alignment file.
- Selections:
 - Cluster size < 5 and no duplicated events.
 - At least three pixel hits and two far apart hits on layer 2.
- Recomputing the cluster positions with GLX2.7 alignment file (Vicente).

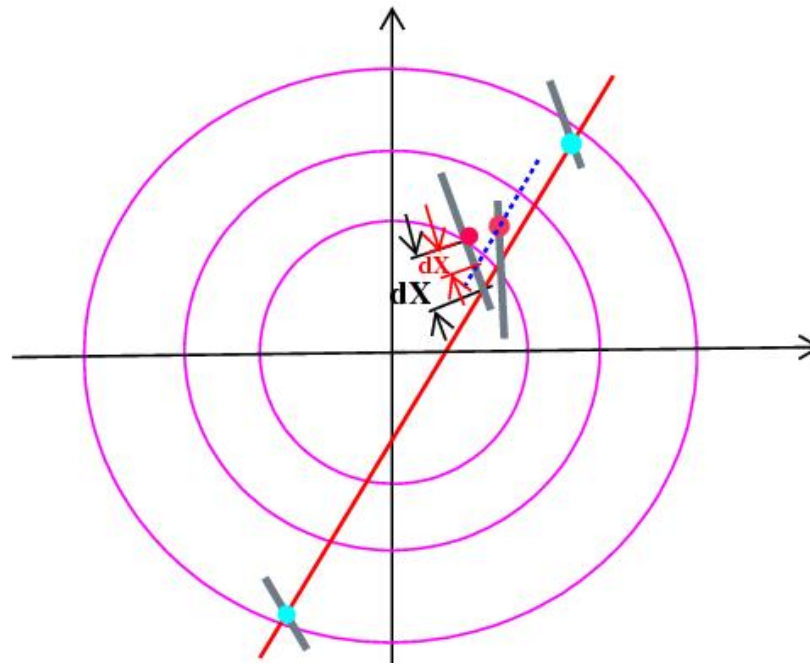


Cosmic Tracks and Overlaps

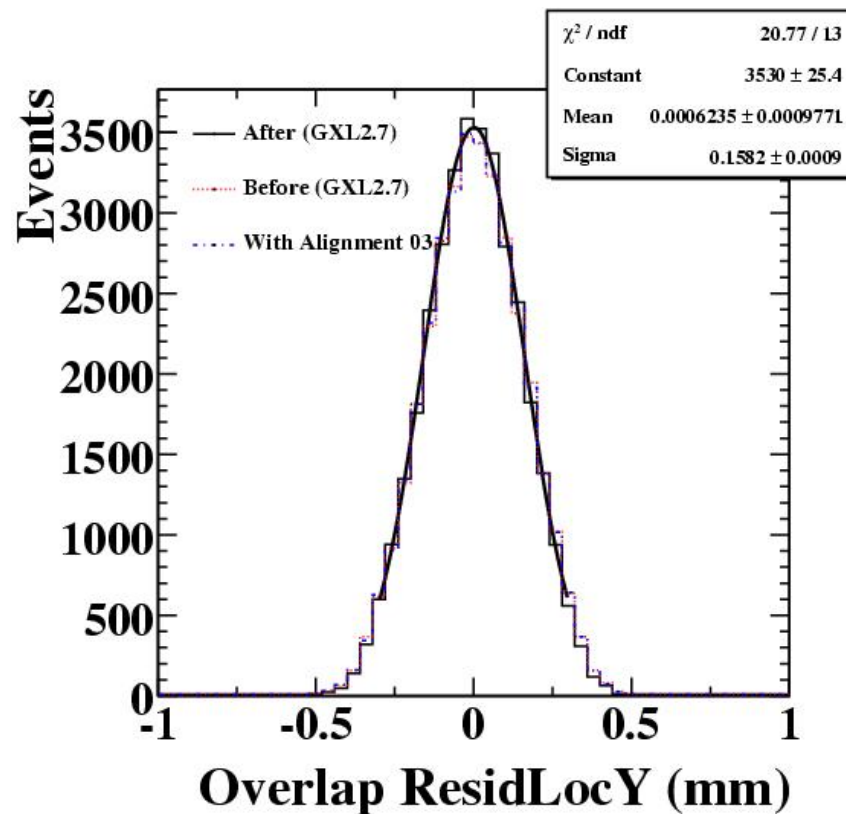
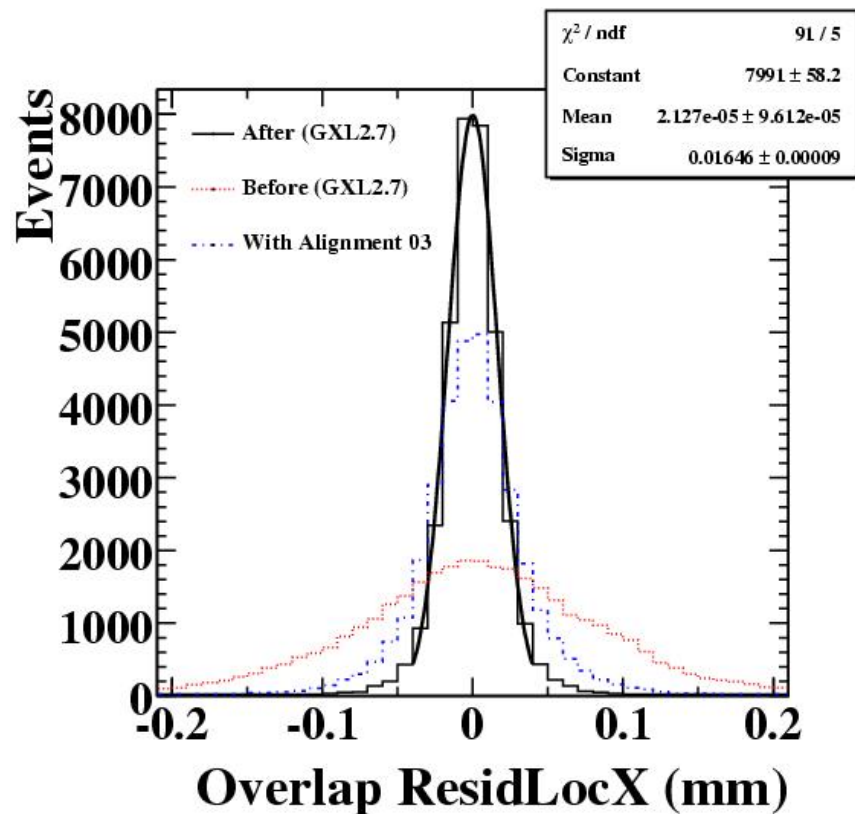


Alignment Strategies

- Step 1: Constraining the relative misalignment between neighboring modules using **overlap residuals**.
- Step 2: Starting two far hits on layer 2 and propagating into layer 0 and 1
- $\chi^2 = \sum (x_{exp} - x_{hit})^2 / \sigma_x^2 + (y_{exp} - y_{hit})^2 / \sigma_y^2$
- There are 86 regions with 5 degrees of freedom each that gives a 430 x 430 matrix to solve in the global χ^2 fit.



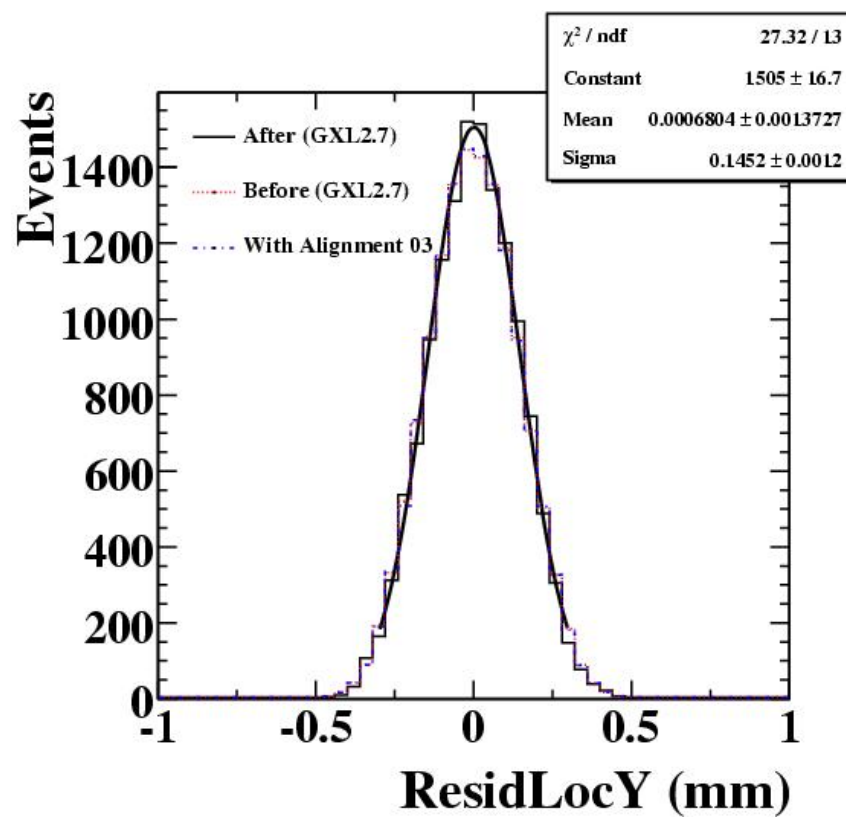
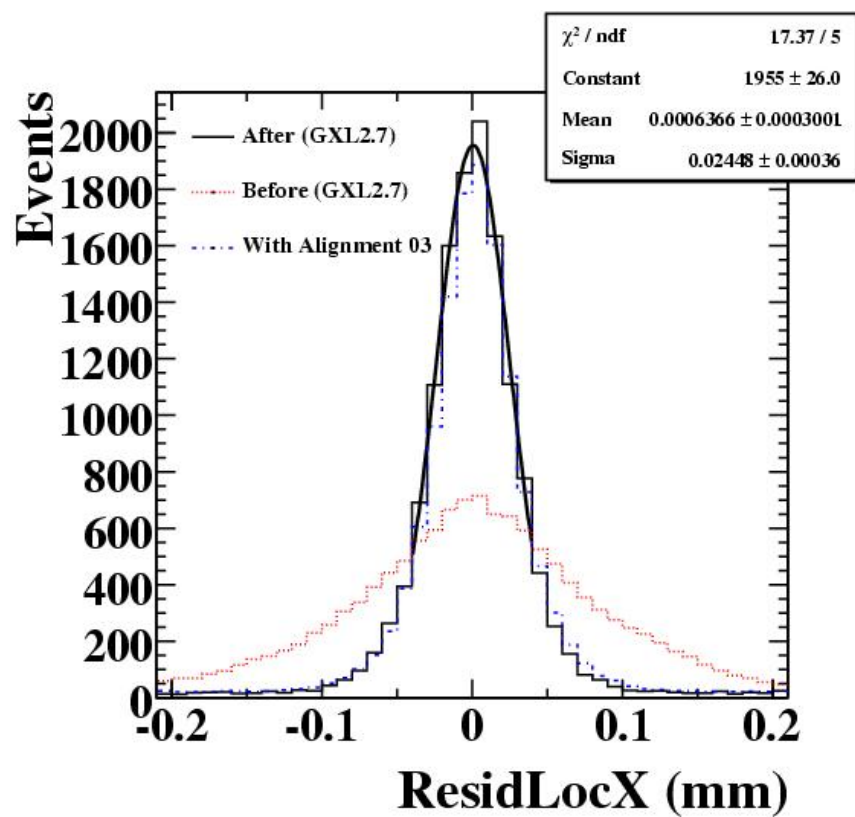
Overlap Residuals after Correction (GXL2.7)



	GXL2.7	Alignment 03	After Correction
Loc σ_X (μm)	77	25.6	16.5
Loc σ_Y (μm)	162	162	158



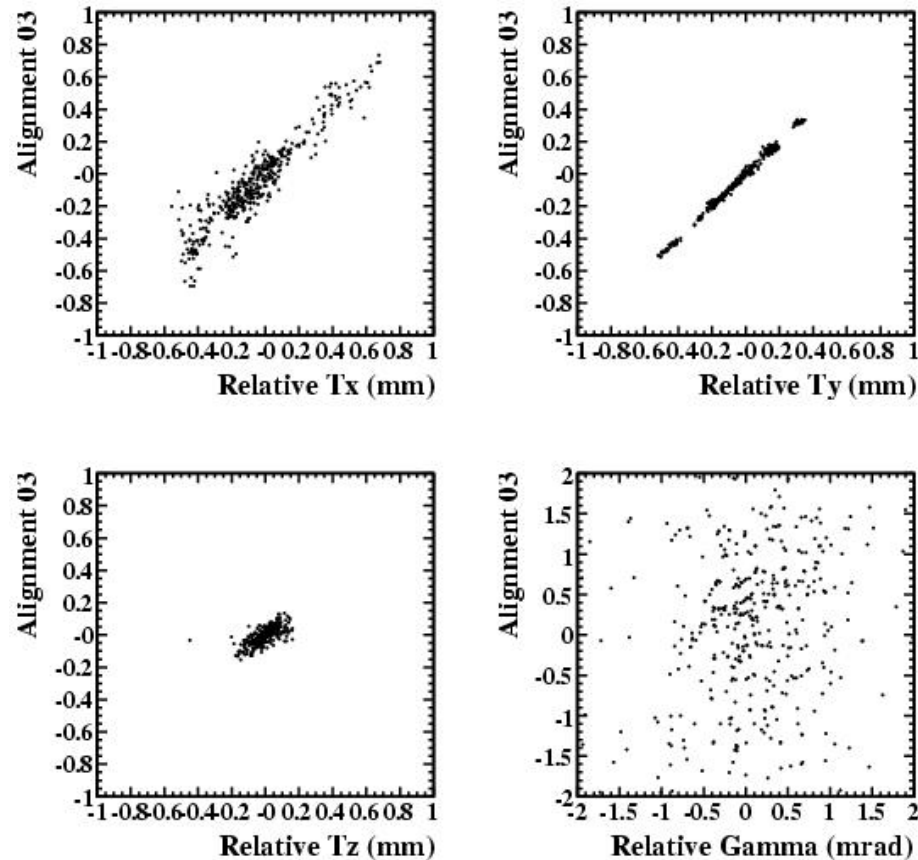
Residuals of layer 0 and 1 after Correction (GXL2.7)



	GXL2.7	Alignment 03	After Correction
Loc σ_X (μm)	73	25.4	24.5
Loc σ_Y (μm)	151	151	145



Comparison of Relative Alignment Constants Between Overlaps



- Derived relative alignment from overlap residuals and compared to alignment 03.
- Consistent with translations, but not much in γ .



Conclusion

- The alternative alignment strategies using overlaps seem promising and give comparable results.
- The overlap residual seem much better, but not much in residuals of layer 0 and 1 (why ?)
- Still statistics limited.
- Interesting to see how the final constants compares to the other methods.

